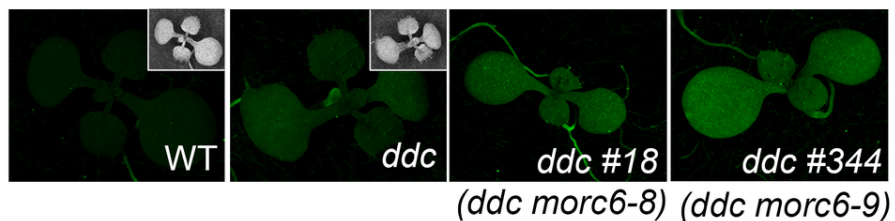
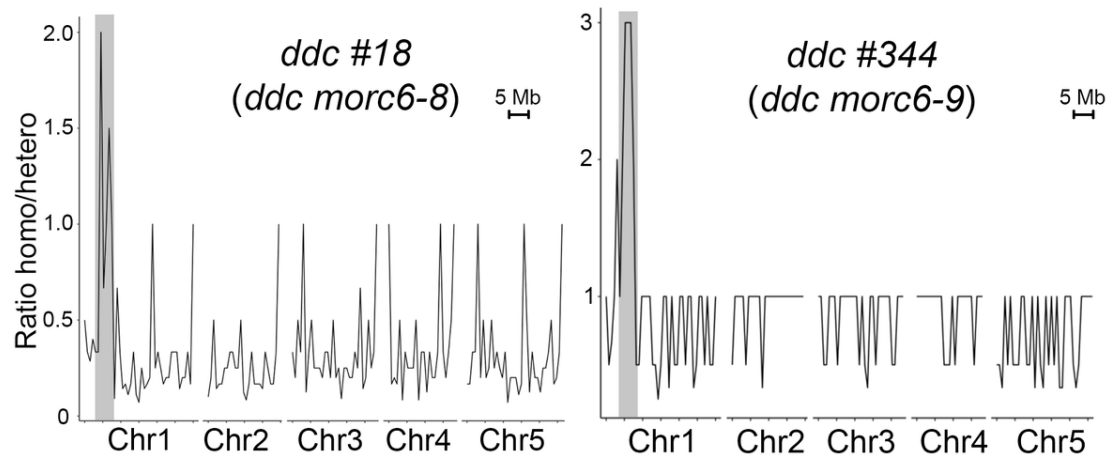


S1 fig

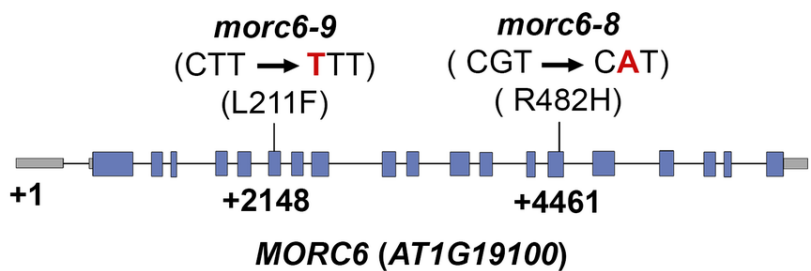
A



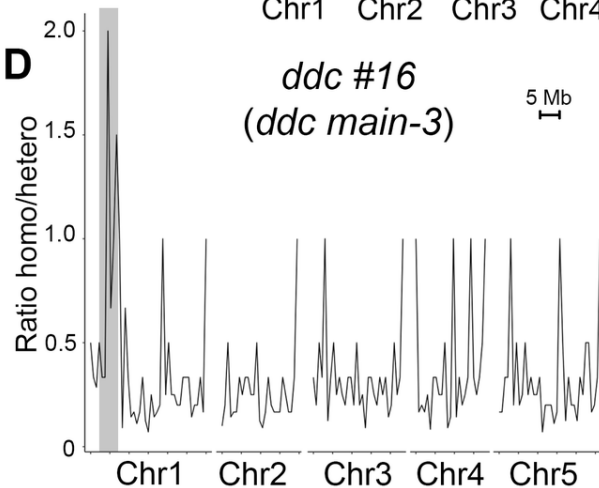
B



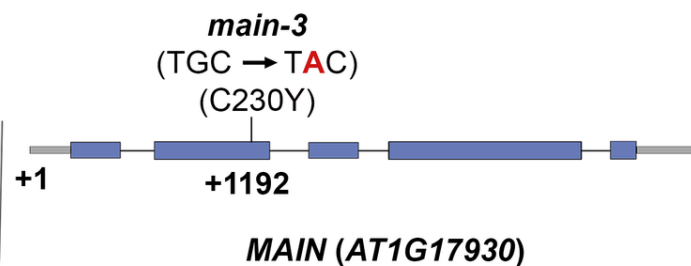
C



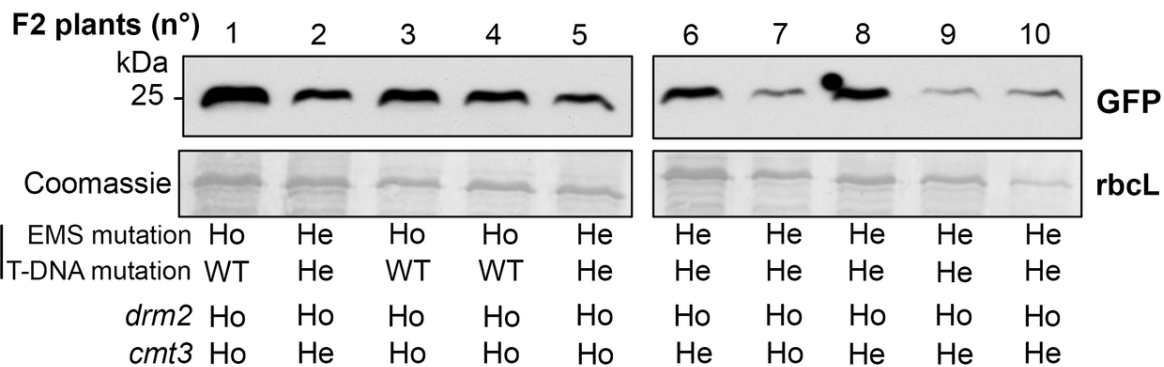
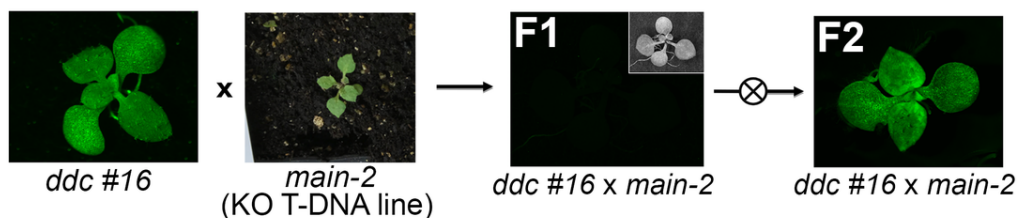
D



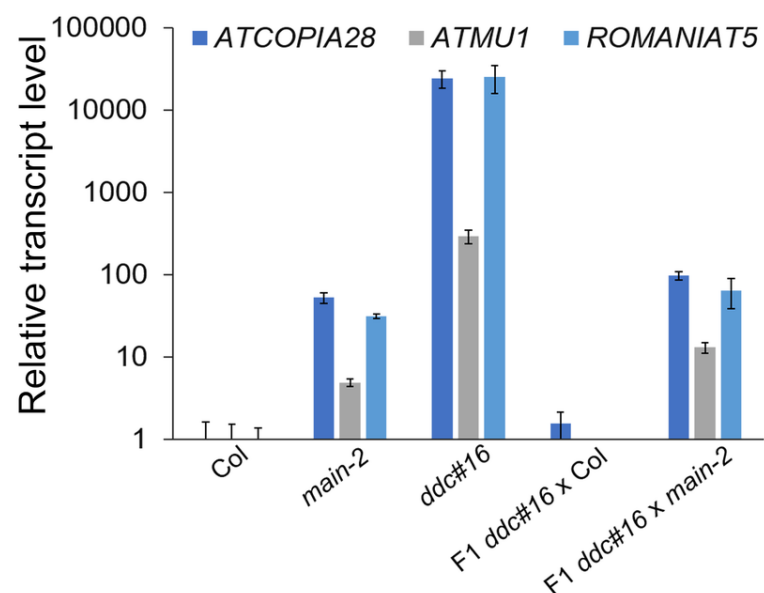
E



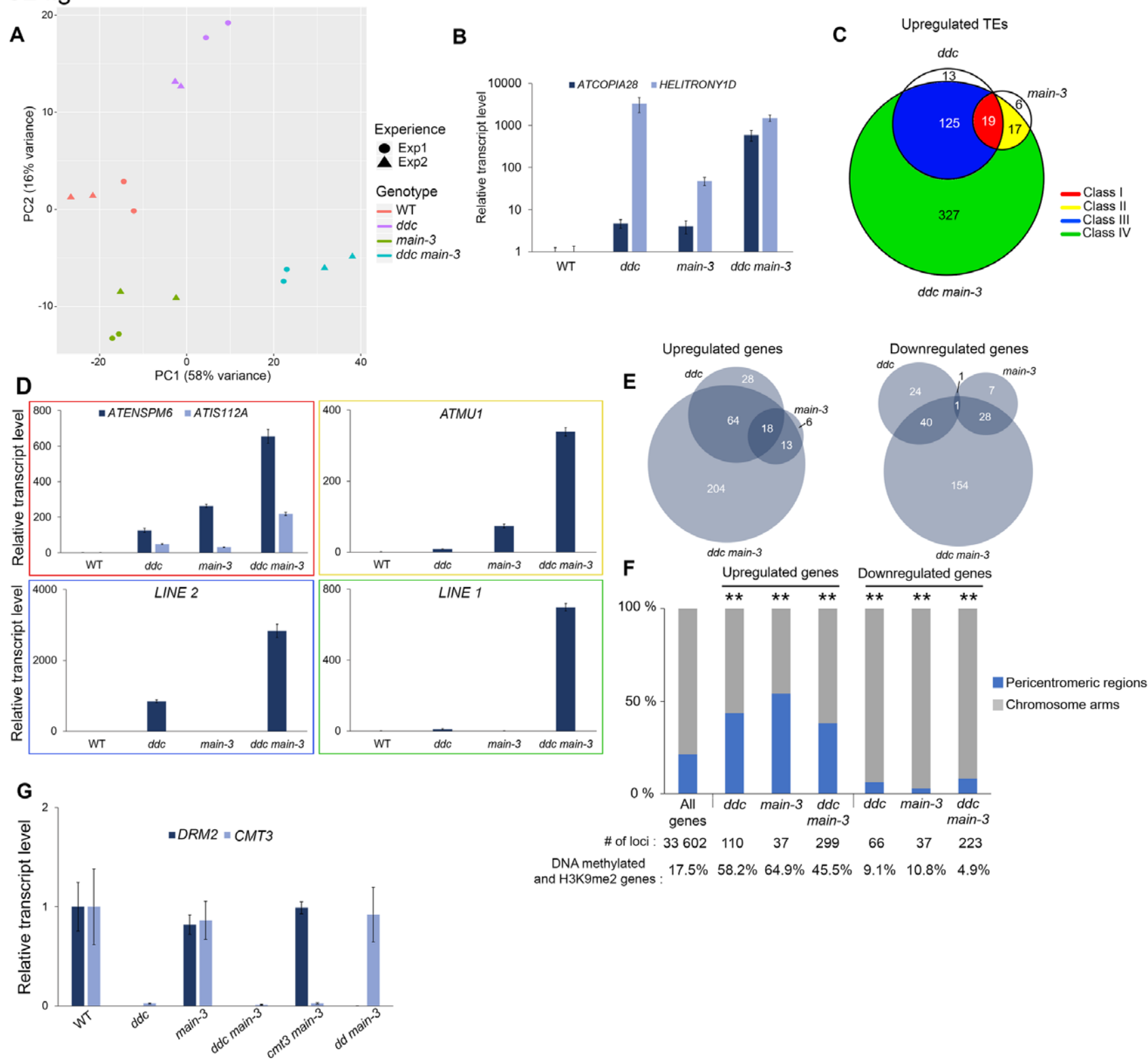
F



G

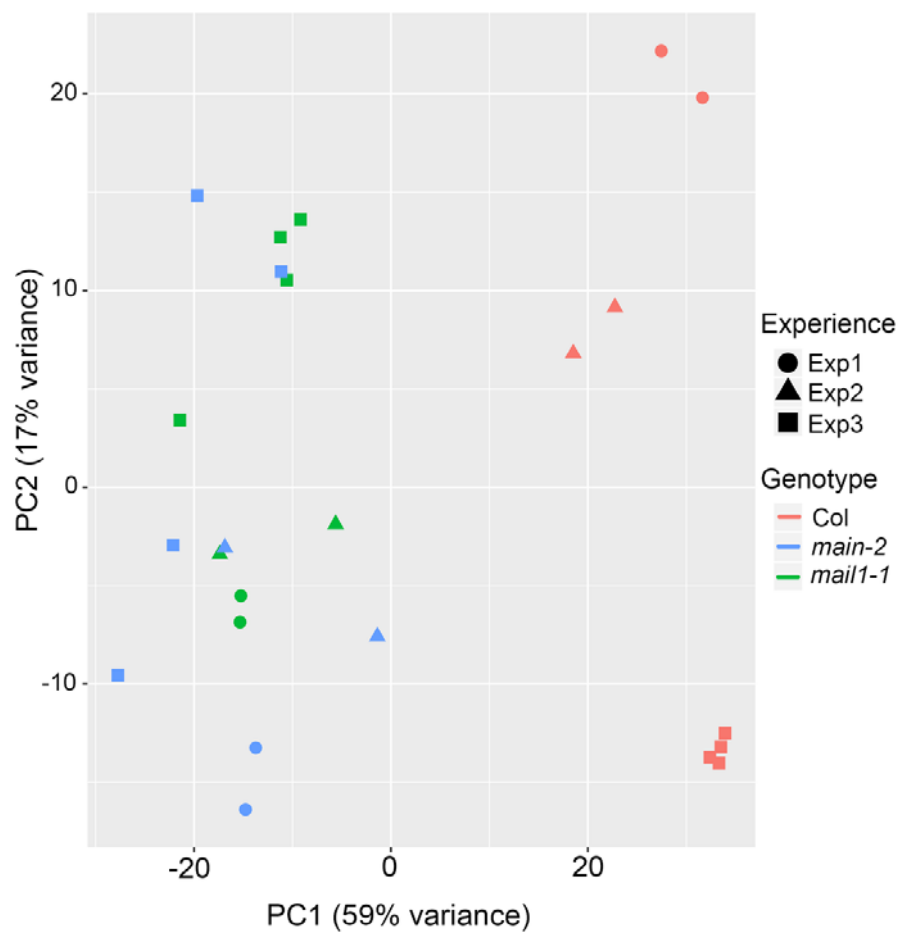


S2 fig

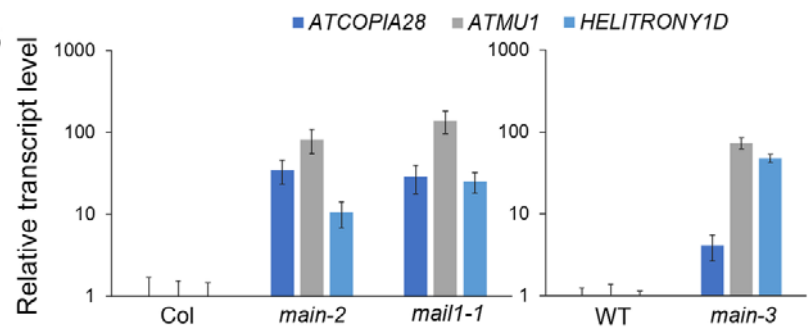


S3 fig

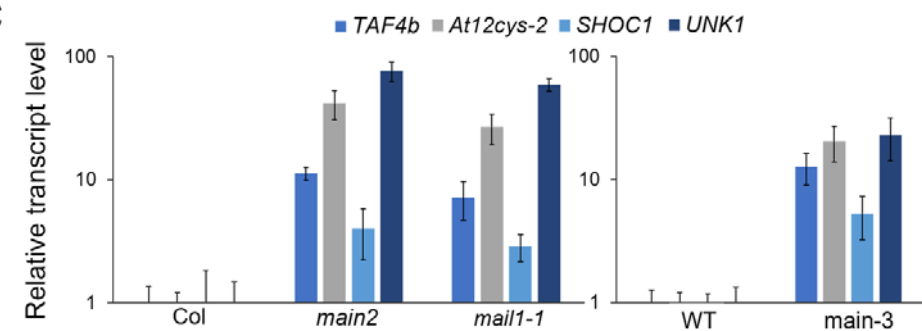
A



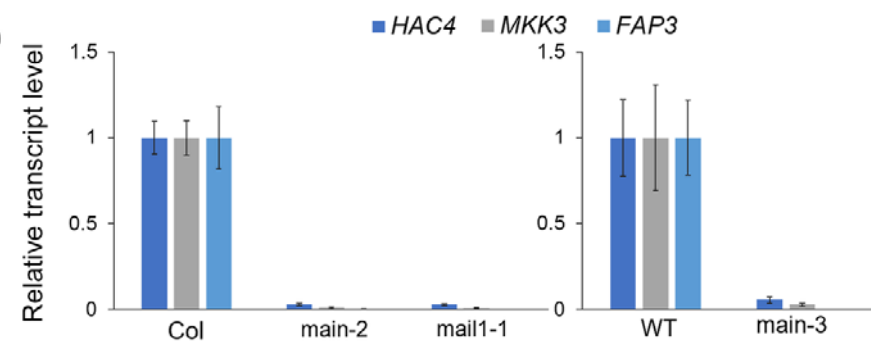
B



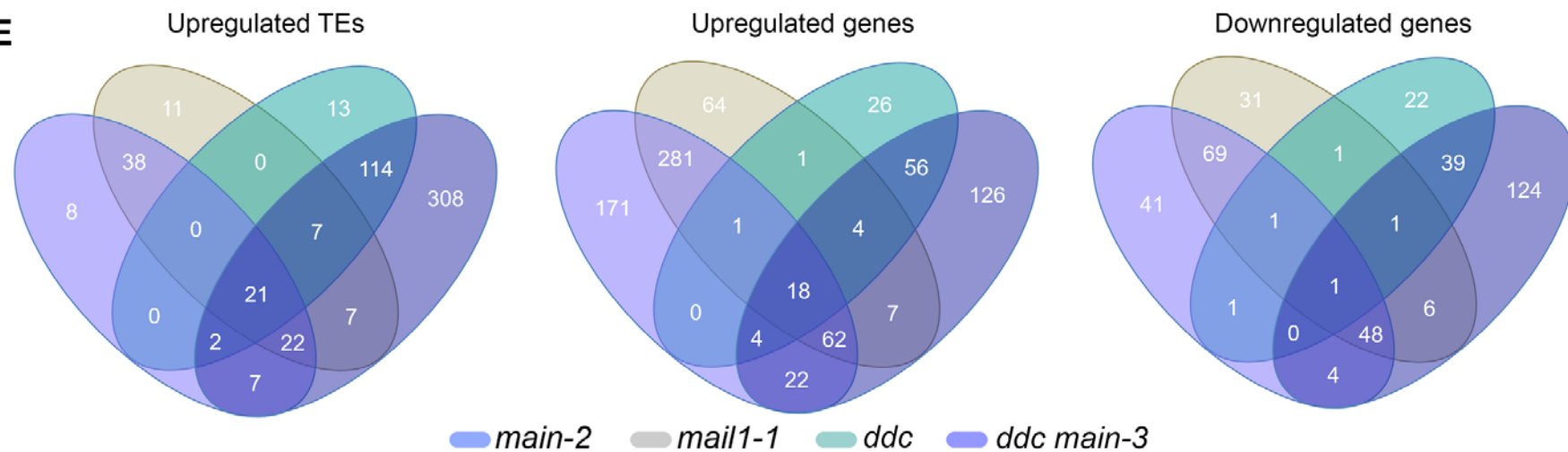
C



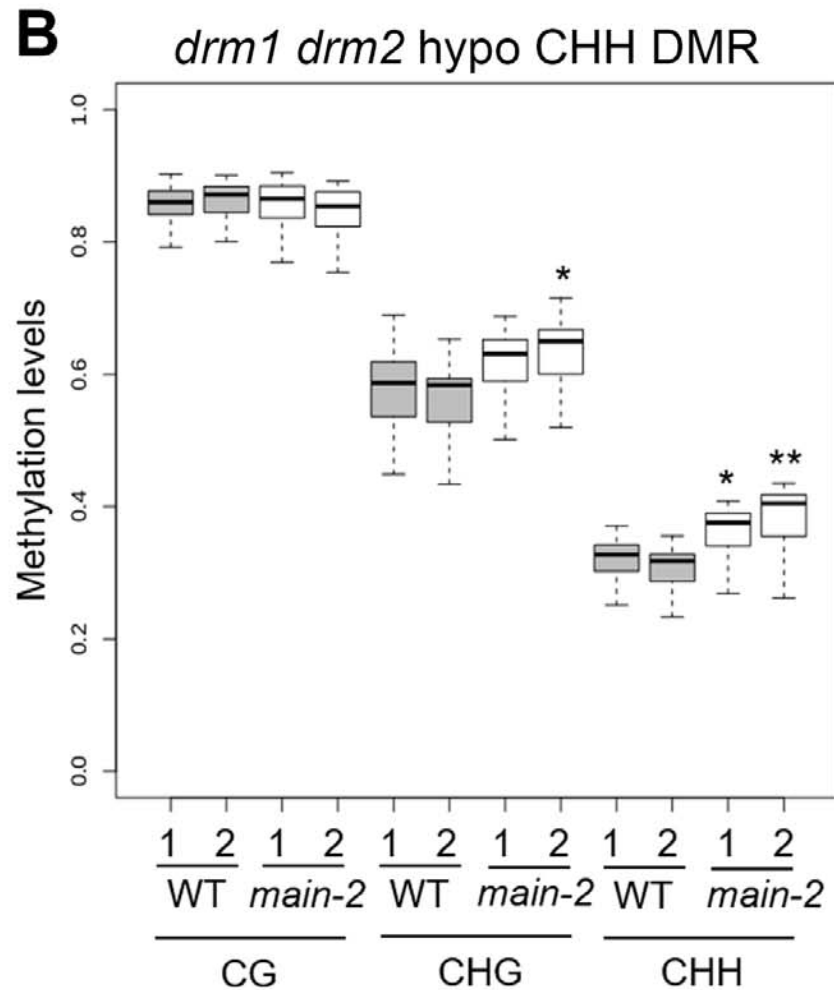
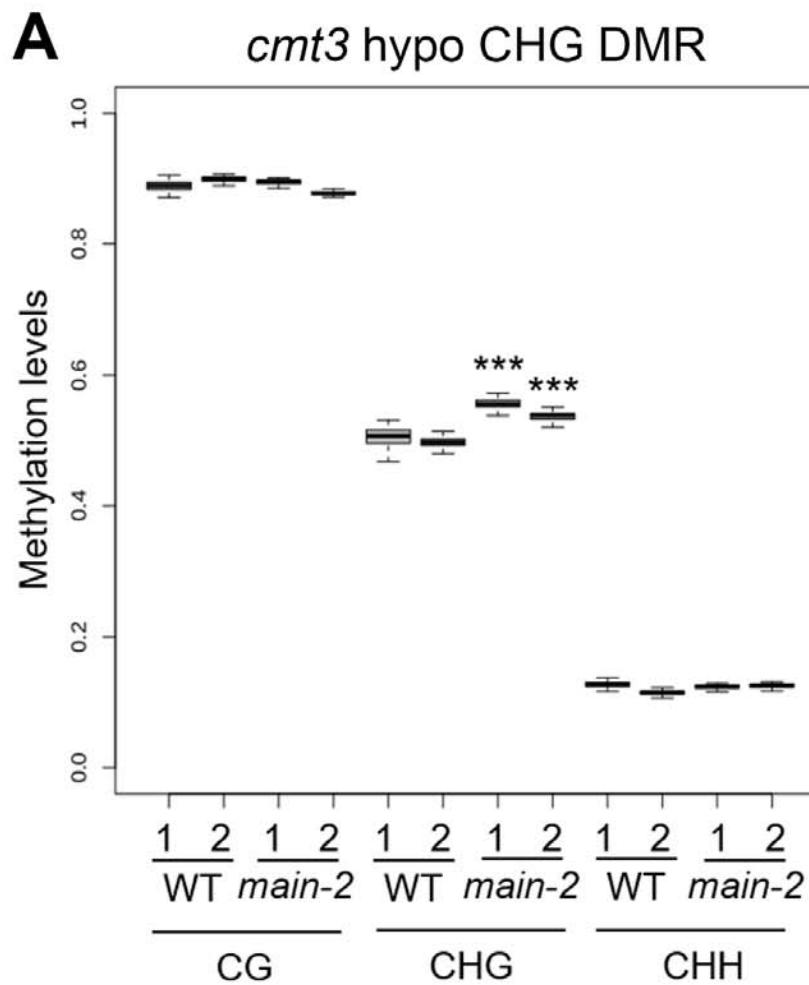
D



E



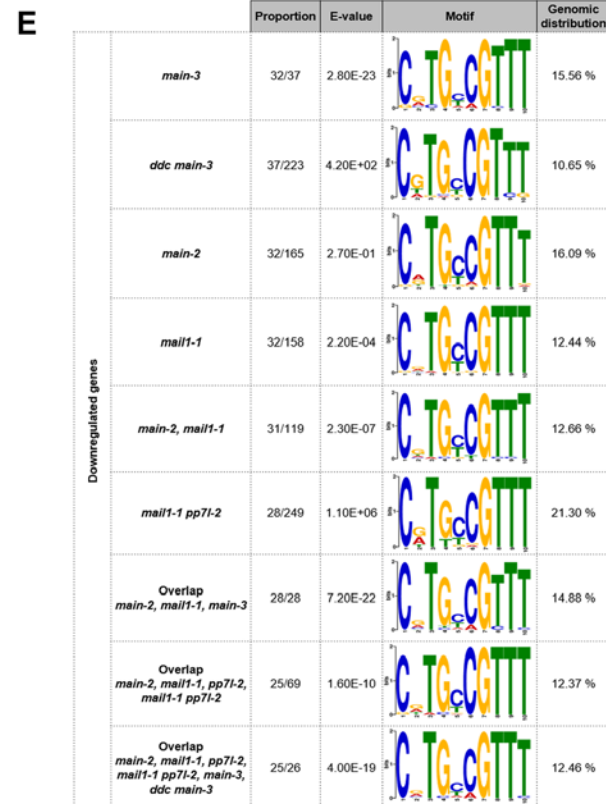
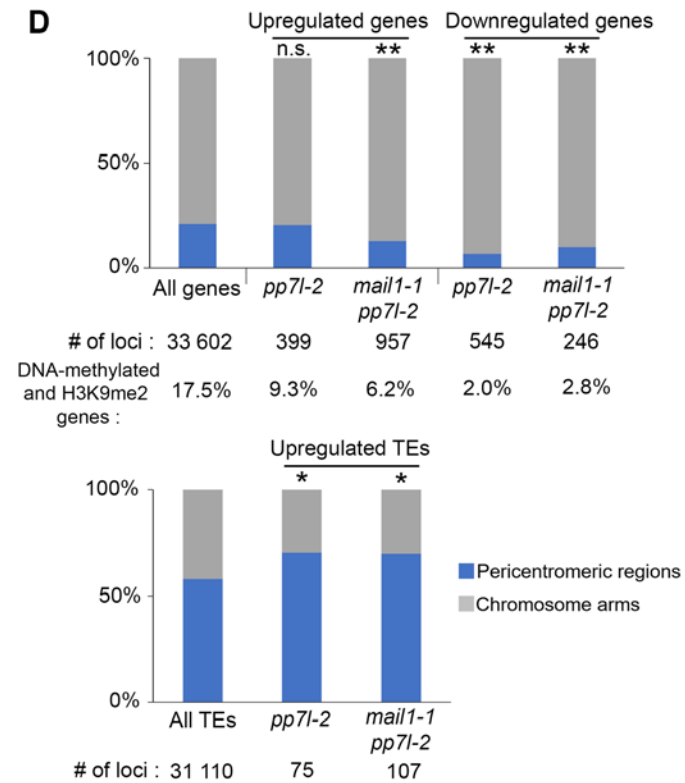
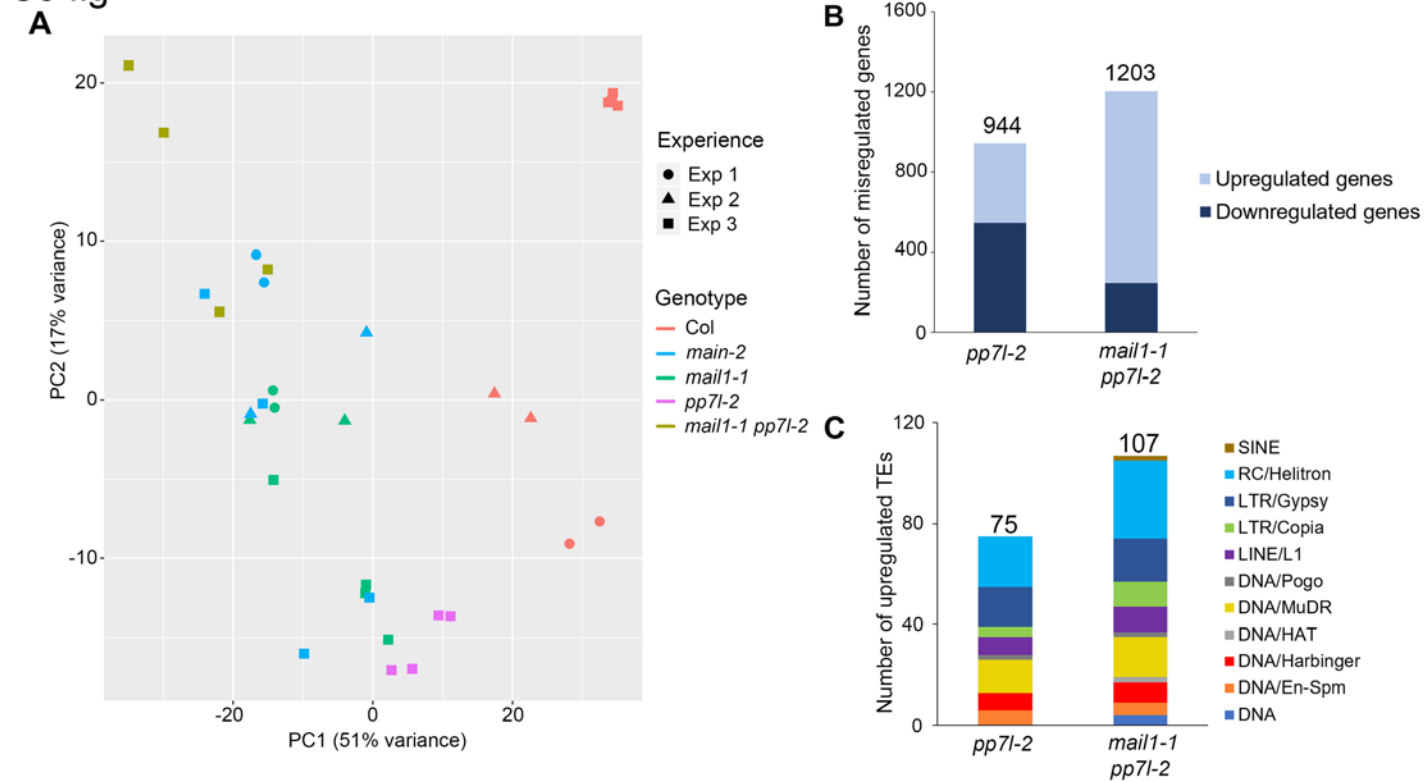
S4 fig



C

		TEs	Genes	Intergenic regions	
<i>main-2</i>	Hyper DRM	CG	4	7	0
		CHG	0	0	1
		CHH	0	0	0
	Hypo DMR	CG	48	28	14
		CHG	0	0	0
		CHH	0	0	0

S5 fig



S6 fig

Fig 1C

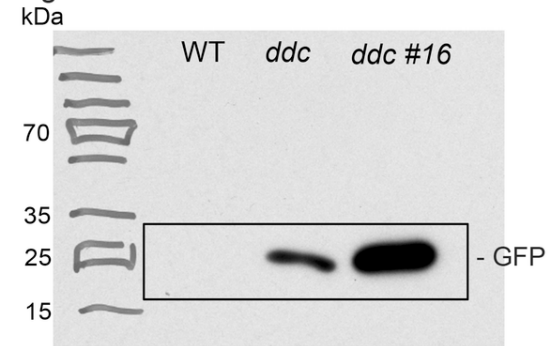


Fig 5G

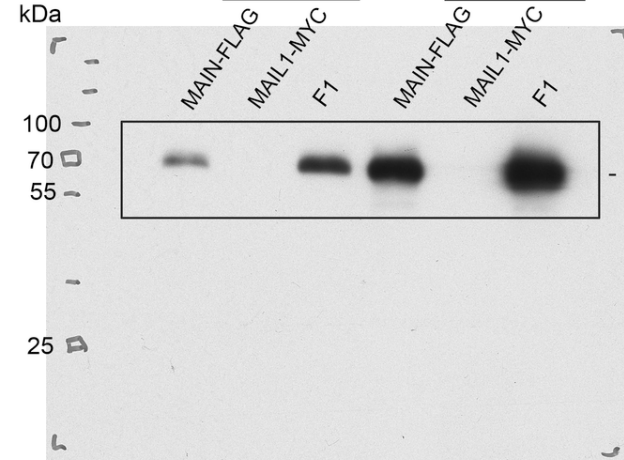


Fig 5G

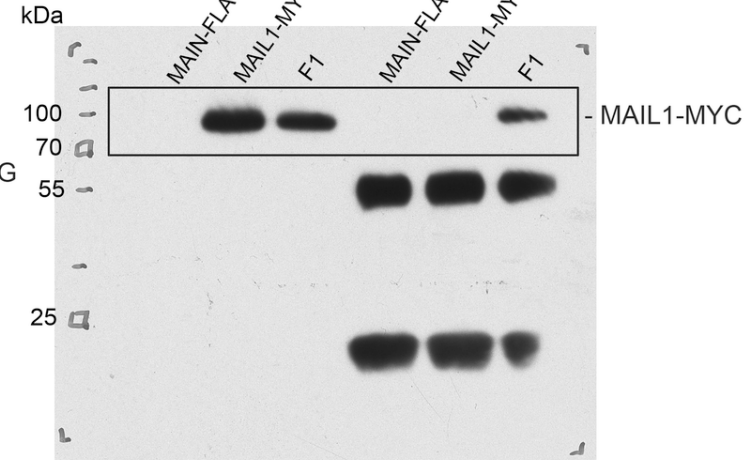


Fig 5B

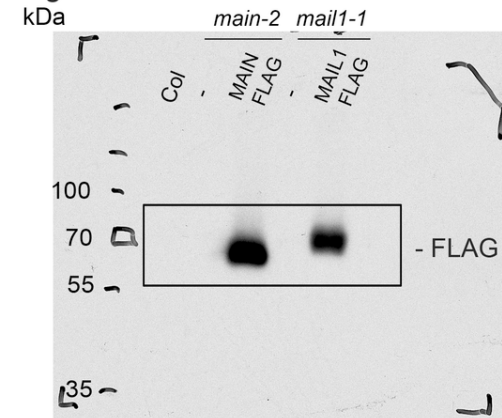


Fig 5H

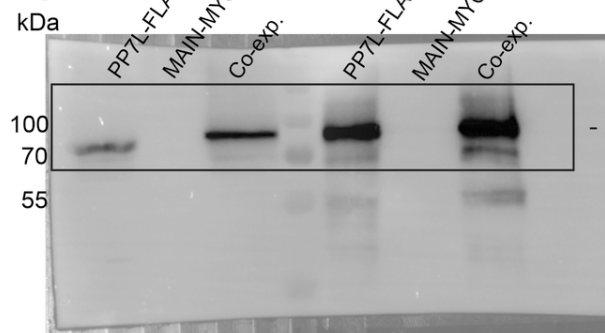


Fig 5H

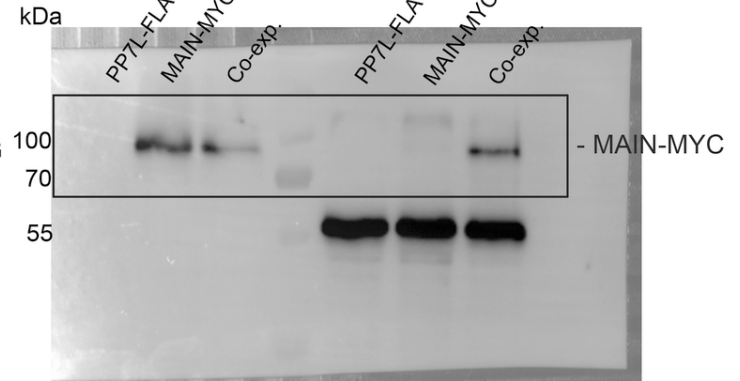


Fig 5B

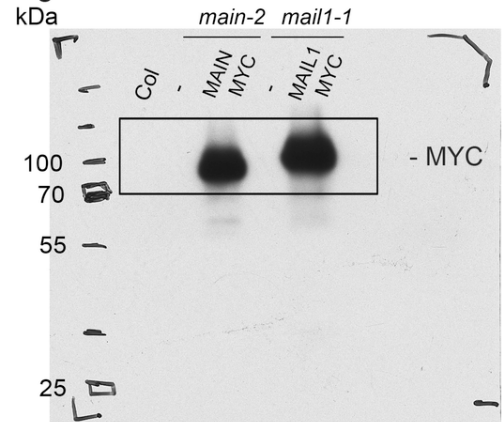


Fig 5I

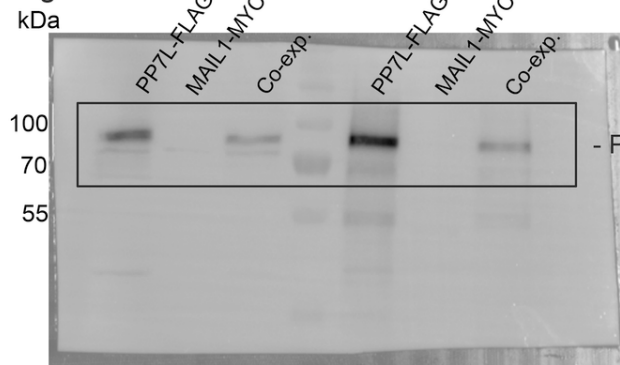


Fig 5I

